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TITLE: Nematode-extracted anticcagulant protein

DATE-ISSUED: February 2, 1999

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US-CL-CURRENT: <u>514/12</u>; <u>530/324</u>, <u>530/350</u>

CLAIMS:

- 1. An isolated protein having Factor VIIa/TF inhibitory activity and having one or more Nematode-extracted Anticoagulant Protein domains ("NAP domains"), wherein each NAP domain includes the sequence;
- Cys-Al-Cys-A2-Cys-A3-Cys-A4-Cys-A5-Cys-A6-Cys-A7-Cys-A8-Cys-A9-Cys-A10, wherein
- (a) Al is an amino acid sequence of 7 to 8 amino acid residues
- (b) Al is an amin: acid sequence;
- c) Ab is an amino acid sequence of 3 amino acid residues;
- (d) A4 is an amin: acid sequence;
- e) At is an amino acid sequence of 3 to 4 amino acid residues;
- f) A6 is an amino acid sequence;
- -g) A7 is an amino abid residue;
- .h) AB is an amino acid sequence of 11 to 12 amino acid residues;
- in Ad is an amino acid sequence of 5 to 7 amino acid residues; and
- j) AlO is an amino acid sequence;
- wherein each of A2, A4, A6 and A10 has an independently selected number of independently selected amino acid residues and each sequence is selected such that each NAF domain has in total less than about 120 amino acid residues and whereir, said isclated protein is derived from a hematophagous nematode species.
- The protein of claim 1, wherein A3 has the sequence Asp-A3.sub.a -A3.sub.b, wherein Al.sub.a and Al.sub.b are independently selected amino acid residues.
- 3. The protein of claim 1, wherein A3 is Asp-Lys-Lys.
- 4. The protein of claim 1, wherein A4 is an amino acid sequence having a net anionis charge.
- 5. The protein of claim 1, wherein A5 has the sequence A5.sub.a -A5.sub.b -A5.sub.c -Ablambld (SE, . II. No. 85), wherein Ablambla through Ablambld are independently selected amino acid residues.
- 6. The protein of plain f, wherein $A\delta$ subta is Leu and $A\delta$ subta is Ara.
- 7. The protein of claim 1, wherein AT is selected from the group consisting of Val and Ile.

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8. The protein of claim 1, wherein Al is Val.
9. The protein of claim 1, wherein Am in ludes an amino acts sequence Amistica
HA8.sub.b HA8.sub.c HA8.sub.i HA8.sub.e HA8.sub.i HA8.sub.i H (CE. II. X.. kej)
wherein
 (a) A8.sub.a is the first amino adid residue in A8,
 (b) at least one of Af.sub.a and Af.sub.b is selected from the group consisting of
Glu or Asp, and
(a) A8.sub.a through A8.sub.g are independently selected amino acid residues.
 10. The protein of claim 9, wherein
(a) A8.sub.a is Glu or Asp,
(b) A8.sub.b is an independently selected amino acid residue,
(c) A8.sub.c is Gly,
(d) A8.sub.d is selected from the group consisting of Fhe, Tyr, and Leu,
(e) A8.sub.e is Tyr,
(f) A6.sub.f is Arg, and
 (g) AS.sub.g is releated from Asp and Asn.
11. The protein of claim 10, wherein -A8.sub.c -A8.sub.d -A8.sub.e -A8.sub.f
-A8.sub.g - is Gly-Phe-Tyr-Arg-Asn [SEQ. ID. NO. 70].
12. The protein of claim 9, wherein
(a) A8.sub.a is an independently selected aming acid residue,
(b) A8.sub.b is Glu or Asp,
(c) A8.sub.c is Gly,
(d) Af.sub.d is selected from the group consisting of Fhe, Tyr, and Leu,
(H) At.sub.e is lyr,
(f) Af.sub.f is Arg, and
(1) Af.sub.q is selected from Asp and Asn.
11. The protein of claim 12, wherein -AB.sub.c -AB.sub.d -AB.sub.e -AB.sub.f
-A8.sub.g - is -Bly-Phe-Tyr-Arg-Ash [SEQ. ID. NO. 70].
14. The protein of claim 1, wherein said nematode species is selected from the group
consisting of Ancylostoma caninum, Ancylostoma ceylanicum, Ancylostoma duodenale,
Necator americanus, and Heligomosomoides polygyrus.
15. The protein of claim 1, wherein
(a) AB is has the sequence Asp-AB.sub.a -AB.sub.p, wherein AB.sub.a and AB.sub.b are
independently selected amino acid residues;
(n) A4 is an amino acid sequence having a net anionic charge;
(r) A5 has the sequence A5.sub.a -A5.sub.b -A5.sub.c -A5.sup.d [SEQ. ID. NO. 85],
wherein A5.sub.a through A5.sub.d are independently selected amino acid residues, and
(d) A7 is selected from the group consisting of Val and Ile.
16. The protein of claim 15 having a NAP domain of AcaNAPc2 (SEQ.\ ID.\ NO.\ 59). 17. The protein of claim 15, wherein said nematode species is selected from the group
consisting of Ancylostoma caninum, Ancylostoma ceylanicum, Ancylostoma duodenale,
Necator americanus, and Heligonesomoides polygyrus.
14. The pritein of plaim 1, wherein
 a) A3 is Asp-Lys-Lys;
(h) A4 is an amino acid sequence having a net anionic charge;
(c) A5 has the sequence A5.sub.a -A5.sub.b -A5.sub.c -A5.sub.d, wherein A5.sub.a is
Leu, A5.sub.c is Ary, and A5.sub.b and A5.sub.d are independently selected amino acid
residues [SEQ. IF. NO. 357],
d) A7 is Val; and
(e) A8 includes an amino acid sequence A3.sub.a -A8.sub.b -Gly -Fhe-Tyr-Arg-Asn [SEQ.
II. NO. 79], wherein at least one of A8.sub.a and A8.sub.b is Glu or Asp.
19. The protein of claim 18 having a NAP domain of AcaNAPc2 (SEQ. ID. NO. 59).
10. The protein of blaim 18, wherein said nematode species is selected from the group
consisting of Ancylostoma caninum, Ancylostoma ceylanicum, Ancylostoma duodenale,
Medator americanus, and Heligomosomoides polygyrus.
11. An isolated protein having Factor VIIa/TF inhibitory activity having a NAP domain
with an amino acid sequence of AcaNAPc2 (SEQ. ID. NO. 59).
11. A pharmaceutical composition comprising the protein of claim 1.
13. A pharmaceutical composition comprising the protein of claim 15.
24. A pharmaceutical composition comprising the protein of claim 19.
25. A pharmaceutical composition comprising an AcaNAPc2 protein [SEQ. ID. NO. 991.
36. A method of inhibiting blood coagulation comprising administering a protein of
claim 1 with a pharmaceutically acceptable carrier.
2\% . A method of inhibiting blood coamplation comprising a ministering a protein \phi :
duaim 15 with a pharmareutically appearance ranner.
28. A method of inhibiting blood obagulation comprising administration against in or
claim 18 with a pharmateutically acceptable carrier.
29. A method of inhibiting blood coagulation comprising a ministering an A WARAC.
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protein (SEQ. ID. NO. E9 .

- 31. A protein of claim 1, wherein said protein has two NAP domains.
- 31. A protein of claim 10, wherein said protein has two MAF domains.
- 32. A protein of claim 1-, wherein said protein has two MAP domains.

unerein said protein is derived from a hematophagous nematode species.

- 33. An isolated protein having anticoagulant activity, wherein said protein specifically inhibits the datalytic activity of the fVIIa. TF complex in the presence
- of fMa or catalytically inactive fMa derivative and not in the absence of fMa or
- catalytically inactive fMa derivative and does not specifically inhibit the activity of fVIIa in the absence of TF and does not specifically inhibit prothrombinase and
- 34. An isolated protein having an amino acid sequence of ASANAPSO SEQ, ID. NO. 59 ,
- wherein said protein specifically inhibits the datalytic activity of the :VIIa TE complex in the presence of fMa or natulytically inactive fMa derivative, and uses not specifically inhibit the activity of EVITA on the ansence of TF and live not
- specifically inhibit prothrombinese.
- 35. An isolated protein having an amino acid sequence of AdaMAF-2 (SE). II. M., 34...
- 36. A protein having an amino acid sequence of AcaMAPc2/proline.
- 37. A method of treating a pathologic condition characterized by abnormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 1.
- 39. A method according to claim 37 wherein said pathologic condition is disseminated intravascular coagulopathy.
- 39. A method of treating a pathologic condition characterized by achormal thrombosis ry preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 15.
- 40. A method according to claim 39 wherein said pathologic condition is disseminated intravascular coaquiopathy.
- 41. A method of treating a pathologic condition characterized by achormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 18.
- 42. A method according to claim 41 wherein said pathologic condition is disseminated intravascular coaquiopathy.
- 43. A method of treating a pathologic condition characterized by aphormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 21.
- 44. A method according to claim 43 wherein said pathologic condition is disseminated intravascular coaquiopathy.
- 45. A method of treating a pathologic condition characterized by appormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 33.
- 46. A method according to claim 45 wherein said pathologic condition is disseminated intravascular coagulopathy.
- 47. A method of treating a pathologic condition characterized by abnormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 35.
- 48. A method according to claim 47 wherein said pathologic condition is disseminated intravascular coadulopathy.
- 49. A method of treating a pathologic condition characterized by abnormal thrombosis by preventing or decreasing said abnormal thrombosis, which comprises administering a protein of claim 36.
- 50. A method according to claim 49 wherein said pathologic condition is disseminated intravascular coaquiopathy.